

**Fig. 1**

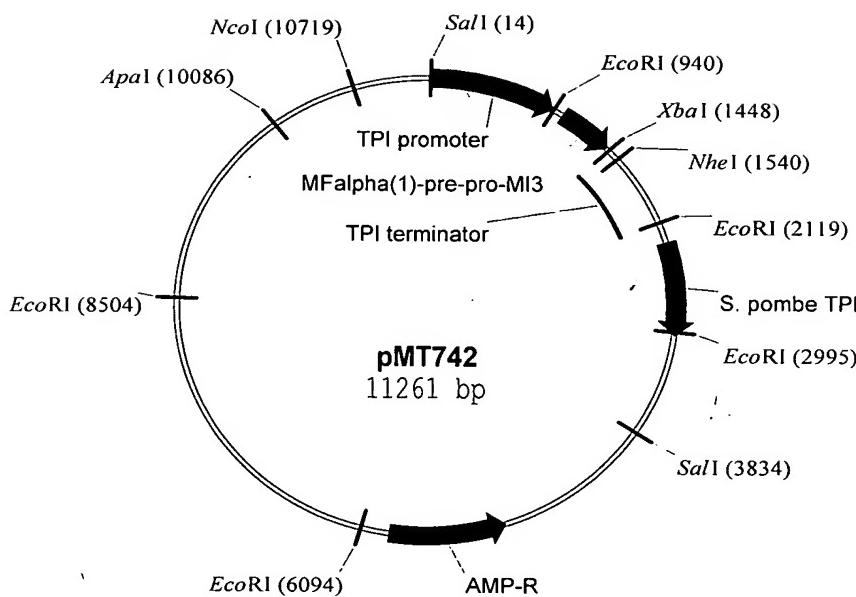


Fig. 2

5

G GA

1 TATAAGAGAG CTCATCTTAT TGTTGTCAGC CCAATGATT CCTTGTCAA  
51 TTGAATTTTC GGATTTACTT GTTCAGGTAC CGCGGTTAAC GGGCTGCCGC  
101 GCCTGTCACT CTAAGAAAAA AGGAGGCCATC AAAAACCCATT CAGCATTAAAC  
10 151 TAAAAACGCG GGTAGAGATT ACTACATATT CCAACAAGAC CTTCGCAGGA  
201 AAGTATAACCT AAACTAATTAA AAGAAATCTC CGAAGTTCGC ATTTCATTGA  
251 ACGGCTCAAT TAATCTTGT AAATATGAGC GTTTTACGT TCACATTGCC  
301 TTTTTTTTA TGTTTACCC TTGCATTTTG GTGCTAAAAG GCGTCACGTT  
351 TTTTTCCGCC GCAGCCGCC CGAAATGAAA AGTATGACCC CCGCTAGACC  
15 401 AAAAATACTT TTGTGTTATT GGAGGATCGC AATCCCTTG GAGCTTTCC  
451 GATACTATCG ACTTATCCGA CCTCTTGTG TTTGAAAATG TCAATTGATA  
501 TCCATCCATT ATATAAAATGC TCAAAACTTG CAGCAACTAT TCTTTACCC  
551 TCCCCGTAA TGGATTGCTA GTCTTAAGGG GGAAATTTGC TGTTTACTAA  
601 AATACAAACC AGGTTTGTG TGGCTTTAT TTGCATTTAA GTAATTACAA  
20 651 TTACAACCAT TAAAAAGAAA ATAAGGCAA ACATATAGCA ATATAATACT  
701 T TC  
701 ATTTACGAAG ATGTCAGCGA TA